RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 09/919,626

Source: /FW16

Date Processed by STIC: 5/1/06

ENTERED

IFW16

RAW SEQUENCE LISTING   DATE: 05/01/2006
PATENT APPLICATION: US/09/919,162G   TIME: 13:11:41

Input Set : A:\PTO.DA.txt
Output Set: N:\CRF4\05012006\I919162G.raw

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      1 <110> APPLICANT: Renauld, Jean-Christophe
      2        Dumoutier, Laure
      4 <120> TITLE OF INVENTION: Isolated Nucleic Acid Molecules Which Encode A Soluble IL-
TIF/IL-22
      5        Receptor or Binding Protein Which Binds to IL-TIF/IL-22, And Uses Thereof
      7 <130> FILE REFERENCE: LUD 5684.2 (10106926)
      9 <140> CURRENT APPLICATION NUMBER: US 09/919,162G
C--> 10 <141> CURRENT FILING DATE: 2001-07-31
     12 <150> PRIOR APPLICATION NUMBER: US 60/245,495
     13 <151> PRIOR FILING DATE: 2000-03-11
     15 <150> PRIOR APPLICATION NUMBER: US 60/234,583
     16 <151> PRIOR FILING DATE: 2000-09-22
     18 <160> NUMBER OF SEQ ID NOS: 11
     20 <210> SEQ ID NO: 1
     21 <211> LENGTH: 21
     22 <212> TYPE: DNA
     23 <213> ORGANISM: Homo sapiens
W--> 24 <400> SEQUENCE: 1
     25 agggtacaat ttcagtcccg a                                    21
     28 <210> SEQ ID NO: 2
     29 <211> LENGTH: 21
     30 <212> TYPE: DNA
     31 <213> ORGANISM: Homo sapiens
W--> 32 <400> SEQUENCE: 2
     33 cggcgtcatg ctccattctg a                                    21
     36 <210> SEQ ID NO: 3
     37 <211> LENGTH: 28
     38 <212> TYPE: DNA
     39 <213> ORGANISM: Homo sapiens
W--> 40 <400> SEQUENCE: 3
     41 tgaacagtca cactcgagac catgatgc                              28
     44 <210> SEQ ID NO: 4
     45 <211> LENGTH: 25
     46 <212> TYPE: DNA
     47 <213> ORGANISM: Homo sapiens
W--> 48 <400> SEQUENCE: 4
     49 catcctgttc tcgaggagct ttaga                                25
     52 <210> SEQ ID NO: 5
     53 <211> LENGTH: 775
     54 <212> TYPE: DNA
     55 <213> ORGANISM: Homo sapiens
W--> 56 <400> SEQUENCE: 5
     57 tgaacagtca cacttgcaac catgatgcct aaacattgct ttctaggctt cctcatcagt   60
     58 ttcttcctta ctggtgtagc aggaactcag tcaacgcatg agtctctgaa gcctcagagg  120
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 59 gtacaatttc agtcccgaaa ttttcacaac attttgcaat ggcagcctgg gagggcactt        180
 60 actggcaaca gcagtgtcta ttttgtgcag tacaaaatat atggacagag acaatggaaa        240
 61 aataaagaag actgttgggg tactcaagaa ctctcttgtg accttaccag tgaaacctca        300
 62 gacatacagg aaccttatta cgggagggtg agggcggcct cggctgggag ctactcagaa        360
 63 tggagcatga cgccgcggtt cactccctgg tgggaaacaa aaatagatcc tccagtcatg        420
 64 aatataaccc aagtcaatgg ctctttgttg gtaattctcc atgctccaaa tttaccatat        480
 65 agataccaaa aggaaaaaaa tgtatctata gaagattact atgaactact ataccgagtt        540
 66 tttataatta caattcact agaaaaggag caaaaggttt atgaaggggc tcacagagcg        600
 67 gttgaaattg aagctctaac accacactcc agctactgtg tagtggctga aatatatcag        660
 68 cccatgttag acagaagaag tcagagaagt gaagagagat gtgtggaaat tccatgactt        720
 69 gtggaatttg gcattcagca atgtggaaat tctaaagctc cctgagaaca ggatg          775
 72 <210> SEQ ID NO: 6
 73 <211> LENGTH: 230
 74 <212> TYPE: PRT
 75 <213> ORGANISM: Homo sapiens
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 77 Met Pro Lys His Cys Phe Leu Gly Phe Leu Ile Ser Phe Phe Leu Thr
 78                   5               10                  15
 79 Gly Val Ala Gly Thr Gln Ser Thr His Glu Ser Leu Lys Pro Gln Arg
 80              20              25              30
 81 Val Gln Phe Gln Ser Arg Asn Phe His Asn Ile Leu Gln Trp Gln Pro
 82          35              40              45
 83 Gly Arg Ala Leu Thr Gly Asn Ser Ser Val Tyr Phe Val Gln Tyr Lys
 84      50              55              60
 85 Ile Tyr Gly Gln Arg Gln Trp Lys Asn Lys Glu Asp Cys Trp Gly Thr
 86 65              70              75              80
 87 Gln Glu Leu Ser Cys Asp Leu Thr Ser Glu Thr Ser Asp Ile Gln Glu
 88              85              90              95
 89 Pro Tyr Tyr Gly Arg Val Arg Ala Ala Ser Ala Gly Ser Tyr Ser Glu
 90              100             105             110
 91 Trp Ser Met Thr Pro Arg Phe Thr Pro Trp Trp Glu Thr Lys Ile Asp
 92          115             120             125
 93 Pro Pro Val Met Asn Ile Thr Gln Val Asn Gly Ser Leu Leu Val Ile
 94      130             135             140
 95 Leu His Ala Pro Asn Leu Pro Tyr Arg Tyr Gln Lys Glu Lys Asn Val
 96 145             150             155             160
 97 Ser Ile Glu Asp Tyr Tyr Glu Leu Leu Tyr Arg Val Phe Ile Ile Asn
 98              165             170             175
 99 Asn Ser Leu Glu Lys Glu Gln Lys Val Tyr Glu Gly Ala His Arg Ala
100              180             185             190
101 Val Glu Ile Glu Ala Leu Thr Pro His Ser Ser Tyr Cys Val Val Ala
102          195             200             205
103 Glu Ile Tyr Gln Pro Met Leu Asp Arg Arg Ser Gln Arg Ser Glu Glu
104      210             215             220
105 Arg Cys Val Glu Ile Pro
106 225             230
109 <210> SEQ ID NO: 7
110 <211> LENGTH: 37
111 <212> TYPE: DNA
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    112 <213> ORGANISM: Homo sapiens
W--> 113 <400> SEQUENCE: 7
    114 ccaacttcca tgatcaatgg aatttccaca catctct                        37
    117 <210> SEQ ID NO: 8
    118 <211> LENGTH: 33
    119 <212> TYPE: DNA
    120 <213> ORGANISM: Homo sapiens
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    122 aagactgagt tgatcaagag aatcgagcct aga                            33
    125 <210> SEQ ID NO: 9
    126 <211> LENGTH: 27
    127 <212> TYPE: DNA
    128 <213> ORGANISM: Homo sapiens
W--> 129 <400> SEQUENCE: 9
    130 aatgtctaga tgctgttctc atttacc                                   27
    133 <210> SEQ ID NO: 10
    134 <211> LENGTH: 792
    135 <212> TYPE: DNA
    136 <213> ORGANISM: Homo sapiens
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    138 atgatgccta acattgctt tctaggcttc ctcatcagtt tcttccttac tggtgtagca    60
    139 ggaactcagt caacgcatga gtctctgaag cctcagaggg tacaatttca gtcccgaaat   120
    140 tttcacaaca tttttgcaatg gcagcctggg agggcactta ctggcaacag cagtgtctat   180
    141 tttgtgcagt acaaaatcat gttctcatgc agcatgaaaa gctctcacca gaagccaagt   240
    142 ggatgctggc agcacatttc ttgtaacttc ccaggctgca gaacattggc taaatatgga   300
    143 cagagacaat ggaaaaataa agaagactgt tggggtactc aagaactctc ttgtgacctt   360
    144 accagtgaaa cctcagacat acaggaacct tattacggga gggtgagggc ggcctcggct   420
    145 gggagctact cagaatggag catgacgccg cggttcactc cctggtggga aacaaaaata   480
    146 gatcctccag tcatgaatat aacccaagtc aatggctctt tgttggtaat tctccatgct   540
    147 ccaaatttac catatagata ccaaaaggaa aaaaatgtat ctatagaaga ttactatgaa   600
    148 ctactatacc gagtttttat aattaacaat tcactagaaa aggagcaaaa ggtttatgaa   660
    149 ggggctcaca gagcggttga aattgaagct ctaacaccac actccagcta ctgtgtagtg   720
    150 gctgaaatat atcagcccat gttagacaga agaagtcaga gaagtgaaga gagatgtgtg   780
    151 gaaattccat ga                                                      792
    154 <210> SEQ ID NO: 11
    155 <211> LENGTH: 263
    156 <212> TYPE: PRT
    157 <213> ORGANISM: Homo sapiens
W--> 158 <400> SEQUENCE: 11
    159 Met Met Pro Lys His Cys Phe Leu Gly Phe Leu Ile Ser Phe Phe Leu
    160                     5                  10                  15
    161 Thr Gly Val Ala Gly Thr Gln Ser Thr His Glu Ser Leu Lys Pro Gln
    162                20                  25                  30
    163 Arg Val Gln Phe Gln Ser Arg Asn Phe His Asn Ile Leu Gln Trp Gln
    164             35                  40                  45
    165 Pro Gly Arg Ala Leu Thr Gly Asn Ser Ser Val Tyr Phe Val Gln Tyr
    166         50                  55                  60
    167 Lys Ile Met Phe Ser Cys Ser Met Lys Ser Ser His Gln Lys Pro Ser
    168 65                  70                  75                  80
```

```
169 Gly Cys Trp Gln His Ile Ser Cys Asn Phe Pro Gly Cys Arg Thr Leu
170              85                  90              95
171 Ala Lys Tyr Gly Gln Arg Gln Trp Lys Asn Lys Glu Asp Cys Trp Gly
172             100             105             110
173 Thr Gln Glu Leu Ser Cys Asp Leu Thr Ser Glu Thr Ser Asp Ile Gln
174         115             120             125
175 Glu Pro Tyr Tyr Gly Arg Val Arg Ala Ala Ser Ala Gly Ser Tyr Ser
176     130             135             140
177 Glu Trp Ser Met Thr Pro Arg Phe Thr Pro Trp Trp Glu Thr Lys Ile
178 145             150             155                 160
179 Asp Pro Pro Val Met Asn Ile Thr Gln Val Asn Gly Ser Leu Leu Val
180             165             170             175
181 Ile Leu His Ala Pro Asn Leu Pro Tyr Arg Tyr Gln Lys Glu Lys Asn
182         180             185             190
183 Val Ser Ile Glu Asp Tyr Tyr Glu Leu Leu Tyr Arg Val Phe Ile Ile
184         195             200             205
185 Asn Asn Ser Leu Glu Lys Glu Gln Lys Val Tyr Glu Gly Ala His Arg
186     210             215             220
187 Ala Val Glu Ile Glu Ala Leu Thr Pro His Ser Ser Tyr Cys Val Val
188 225             230             235                 240
189 Ala Glu Ile Tyr Gln Pro Met Leu Asp Arg Arg Ser Gln Arg Ser Glu
190             245             250             255
191 Glu Arg Cys Val Glu Ile Pro
192             260
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VERIFICATION SUMMARY   DATE: 05/01/2006
PATENT APPLICATION: **US/09/919,162G**   TIME: 13:11:42

Input Set : **A:\PTO.DA.txt**
Output Set: **N:\CRF4\05012006\I919162G.raw**

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L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:24 M:283 W: Missing Blank Line separator, <400> field identifier
L:32 M:283 W: Missing Blank Line separator, <400> field identifier
L:40 M:283 W: Missing Blank Line separator, <400> field identifier
L:48 M:283 W: Missing Blank Line separator, <400> field identifier
L:56 M:283 W: Missing Blank Line separator, <400> field identifier
L:76 M:283 W: Missing Blank Line separator, <400> field identifier
L:113 M:283 W: Missing Blank Line separator, <400> field identifier
L:121 M:283 W: Missing Blank Line separator, <400> field identifier
L:129 M:283 W: Missing Blank Line separator, <400> field identifier
L:137 M:283 W: Missing Blank Line separator, <400> field identifier
L:158 M:283 W: Missing Blank Line separator, <400> field identifier
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